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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/509,945

DATE: 07/30/2001
TIME: 15:59:23

Input Set : A:\0230-0148P.ST25.txt
Output Set: N:\CRF3\07302001\I509945.raw

3 <110> APPLICANT: HAMADA, Kazuyuki et al.
5 <120> TITLE OF INVENTION: MUTANT BARNASE GENE AND TRANSGENIC PLANT TRANSFORMED BY SAID
GENE
7 <130> FILE REFERENCE: 0230-0148P
9 <140> CURRENT APPLICATION NUMBER: 09/509,945
C--> 10 <141> CURRENT FILING DATE: 2001-04-01
12 <160> NUMBER OF SEQ ID NOS: 7
14 <170> SOFTWARE: PatentIn version 3.1
16 <210> SEQ ID NO: 1
17 <211> LENGTH: 343
18 <212> TYPE: DNA
19 <213> ORGANISM: *Bacillus amyloliquefaciens*
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22 <221> NAME/KEY: misc_feature
23 <223> OTHER INFORMATION: wild type barnase gene
26 <220> FEATURE:
27 <221> NAME/KEY: CDS
28 <222> LOCATION: (1)..(336)
29 <223> OTHER INFORMATION:
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35 1 5 10 15
37 aca tat cat aag cta cct gat aat tac att aca aaa tca gaa gca caa 96
38 Thr Tyr His Lys Leu Pro Asp Asn Tyr Ile Thr Lys Ser Glu Ala Gln
39 20 25 30
41 gcc ctc ggc tgg gtg gca tca aaa ggg aac ctt gca gac gtc gct ccg 144
42 Ala Leu Gly Trp Val Ala Ser Lys Gly Asn Leu Ala Asp Val Ala Pro
43 35 40 45
45 ggg aaa agc atc ggc gga gac atc ttc tca aac agg gaa ggc aaa ctc 192
46 Gly Lys Ser Ile Gly Asp Ile Phe Ser Asn Arg Glu Gly Lys Leu
47 50 55 60
49 ccg ggc aaa agc gga cga aca tgg cgt gaa gcg gat att aac tat aca 240
50 Pro Gly Lys Ser Gly Arg Thr Trp Arg Glu Ala Asp Ile Asn Tyr Thr
51 65 70 75 80
53 tca ggc ttc aga aat tca gac cgg att ctt tac tca agc gac tgg ctg 288
54 Ser Gly Phe Arg Asn Ser Asp Arg Ile Leu Tyr Ser Ser Asp Trp Leu
55 85 90 95
57 att tac aaa aca acg gac cat tat cag acc ttt aca aaa atc aga taa 336
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61 ggtaacc 343
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65 <211> LENGTH: 111
66 <212> TYPE: PRT
67 <213> ORGANISM: *Bacillus amyloliquefaciens*
69 <220> FEATURE:
70 <221> NAME/KEY: misc_feature

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83 Ala Leu Gly Trp Val Ala Ser Lys Gly Asn Leu Ala Asp Val Ala Pro
84 35 40 45
87 Gly Lys Ser Ile Gly Gly Asp Ile Phe Ser Asn Arg Glu Gly Lys Leu
88 50 55 60
91 Pro Gly Lys Ser Gly Arg Thr Trp Arg Glu Ala Asp Ile Asn Tyr Thr
92 65 70 75 80
95 Ser Gly Phe Arg Asn Ser Asp Arg Ile Leu Tyr Ser Ser Asp Trp Leu
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105 <212> TYPE: DNA
106 <213> ORGANISM: Artificial Sequence
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116 agggAACCTT gcagacgtcg ctccggggaa aagcatcgcc ggagacatct tctcaaacag 180
118 ggaaggcaaa ctccggggca aaagcggacg aacatggcgt gaagcggata ttaactatac 240
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142 ccctttttg cggcattttg cttccctgtt tttgctcacc cagaaacgct ggtgaaagta 240
144 aaagatgtg aagatcgatgggtgcacga gtgggttaca tcgaactggc tctcaacagc 300
146 ggtaagatcc ttgagagttt tcgccccgaa gaacgttttc caatgtatgg cactttaaa 360
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156 aacatggggg atcatgtaac tcgccttgat cgttggaaac cggagctgaa tgaagccata 660
158 ccaaacgacg agcgtgacac cacgatgcct gtagcaatgg caacaacgtt gcgcaaaacta 720

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166	aagccctcccc gtatcgtagt tatctacacg acggggagtc aggcaactat ggatgaacga	960
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170	gtttactcat atatacttta gattgattta aaacttcatt ttaattttaa aaggatctag	1080
172	gtgaagatcc tttttggctc gagtctcatg accaaaatcc cttAACGTA gtttgcgttc	1140
174	cactgagcgt cagaccccggt agaaaagatc aaaggatctt cttgagatcc ttttttctg	1200
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